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OM nucleic - nucleic search, using sw model

Run on: March 30, 2003, 03:17:01 ; Search time 85.7612 Seconds
(without alignments)
13395.488 Million cell updates/sec

Title: US-09-768-781-1
Perfect score: 1350
Sequence: 1 atggacagagtttatgaat.....caaggcaaaagtgtgtctga 1350

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 574371 seqs, 425486471 residues

Total number of hits satisfying chosen parameters: 1148742

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Published Applications NA:*

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- 10: /cgn2_6/ptodata/1/pubpna/US09_PUBCOMB.seq:*
- 11: /cgn2_6/ptodata/1/pubpna/US10_NEW_PUB.seq:*
- 12: /cgn2_6/ptodata/1/pubpna/US10_PUBCOMB.seq:*
- 13: /cgn2_6/ptodata/1/pubpna/US60_NEW_PUB.seq:*
- 14: /cgn2_6/ptodata/1/pubpna/US60_PUBCOMB.seq:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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2	1346.8	99.8	1389	10	US-09-768-781-2
3	747.2	55.3	17993	10	US-09-768-781-5
4	513.8	38.1	531	9	US-10-092-154-106
5	513.8	38.1	531	10	US-09-764-847-106
6	289.6	21.5	5096	10	US-09-962-436-564
7	270.6	20.0	668	10	US-09-864-761-16902
8	176.8	13.1	471	10	US-09-864-761-62
9	160.4	11.9	384	10	US-09-864-761-21423
10	140.4	10.4	498	10	US-09-864-761-4683
11	73.2	5.4	294	10	US-09-864-761-19197
12	65	4.8	477	10	US-09-864-761-2467
13	51.6	3.8	486	10	US-09-864-761-10062
14	43.4	3.2	832	10	US-09-764-877-853
15	38.8	2.9	6799	9	US-09-902-941-1883
16	38.8	2.9	6799	9	US-09-849-626-1883
17	38.8	2.9	6799	9	US-10-017-754-1883
18	36.4	2.7	3504	9	US-09-822-846-143
19	36.4	2.7	3504	9	US-09-822-846-144

20	34.8	2.6	2120	10	US-09-801-574-29	Sequence 29, Appl
c 21	34.4	2.5	42000	9	US-10-081-563-25	Sequence 25, Appl
c 22	34.4	2.5	465237	10	US-09-933-267A-1	Sequence 1, Appl
c 23	34	2.5	170834	10	US-09-835-232-7	Sequence 7, Appl
c 24	34	2.5	536165	9	US-09-939-964-1	Sequence 1, Appl
c 25	33.6	2.5	9822	10	US-09-853-386-25	Sequence 25, Appl
c 26	33.6	2.5	22756	9	US-10-091-572-473	Sequence 473, Appl
c 27	33.4	2.5	334	10	US-09-867-701-9820	Sequence 9820, Ap
c 28	33.4	2.5	592	10	US-09-864-761-13139	Sequence 13139, A
c 29	33.4	2.5	1233	9	US-10-076-816-12	Sequence 12, Appl
c 30	33.4	2.5	1940	12	US-10-044-090-275	Sequence 275, App
c 31	33.4	2.5	11337	10	US-09-764-877-2651	Sequence 2651, Ap
c 32	33.2	2.5	327	10	US-09-864-761-28059	Sequence 28059, A
c 33	33.2	2.5	456	10	US-09-864-761-11468	Sequence 11468, A
c 34	33.2	2.5	4030	10	US-09-070-927A-264	Sequence 264, App
c 35	33	2.4	1148	10	US-09-893-737-35	Sequence 35, Appl
c 36	33	2.4	4689	10	US-09-895-652-10	Sequence 10, Appl
c 37	33	2.4	5173	10	US-09-811-045A-2	Sequence 2, Appl
c 38	33	2.4	1503841	9	US-09-946-807-1	Sequence 1, Appl
c 39	33	2.4	1503841	10	US-09-795-668-1	Sequence 1, Appl
c 40	33	2.4	1503841	10	US-09-795-686-1	Sequence 1, Appl
c 41	32.8	2.4	817	10	US-09-864-761-18944	Sequence 18944, A
c 42	32.8	2.4	953	10	US-09-864-761-2203	Sequence 2203, Ap
c 43	32.8	2.4	2125	9	US-09-957-708-19	Sequence 19, Appl
c 44	32.8	2.4	5598	9	US-09-938-842A-1436	Sequence 1436, Ap
c 45	32.6	2.4	222	10	US-09-915-060-1	Sequence 1, Appl

ALIGNMENTS

RESULT 1
US-09-768-781-1
; Sequence 1, Application US/09768781
; Patent No. US20020142376A1
; GENERAL INFORMATION:
; APPLICANT: MERKULOV, Gennady V. et al
; TITLE OF INVENTION: ISOLATED HUMAN TRANSPORTER PROTEINS,
; TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING HUMAN TRANSPORTER PROTEINS,
; FILE REFERENCE: CL001057-CIP
; CURRENT APPLICATION NUMBER: US/09/768,781
; CURRENT FILING DATE: 2001-01-25
; NUMBER OF SEQ ID NOS: 7
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 1
; LENGTH: 1350
; TYPE: DNA
; ORGANISM: Human
; US-09-768-781-1

Query Match	100.0%;	Score 1350;	DB 10;	Length 1350;
Best Local Similarity	100.0%;	Pred. No. 0;		
Matches 1350;	Conservative 0;	Mismatches 0;	Indels 0;	Gaps 0;
Qy	1	ATGGACAGAGTTTATGAATTCCTGAGGAGCCAAATGTGGATCCGGTTTCATCTCTGGAG	60	
Db	1	ATGGACAGAGTTTATGAATTCCTGAGGAGCCAAATGTGGATCCGGTTTCATCTCTGGAG	60	
Qy	61	GAAGATGTCATCCGTGGAGCCAAACCCCGATTACTTTTCCATTAGCATCTTTTCTCC	120	
Db	61	GAAGATGTCATCCGTGGAGCCAAACCCCGATTACTTTTCCATTAGCATCTTTTCTCC	120	
Qy	121	ACCTTTTGTACTGTGGGAGGCTGTCCTTTGTACATGTTAGAAATCTATCGAAG	180	
Db	121	ACCTTTTGTACTGTGGGAGGCTGTCCTTTGTACATGTTAGAAATCTATCGAAG	180	
Qy	181	AATAGTGAATCTACCGGATGACATACACCTTTTCTTTTATGTTTTCATCCATTATG	240	
Db	181	AATAGTGAATCTACCGGATGACATACACCTTTTCTTTTATGTTTTCATCCATTATG	240	
Qy	241	GTCAGTTGACCTTCATTTTGTGCCAGAGATCTAGCCAAAGATAAACCGCTATCATTA	300	


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QY 721 CTCTGCATCACCATCGCGGACATTGGAGATCACTTCCCGCTCCTGATTCCTGGTGCTC 780
DB 760 CTCTGCATCACCATCGCGGACATTGGAGATCACTTCCCGCTCCTGATTCCTGGTGCTC 819
QY 781 TTCTCAGCCACTTGAATTAAGAGTGTGCGCTTCTAGTGTCTCAACTTCTGTATCATC 840
DB 820 TTCTCAGCCACTTGAATTAAGAGTGTGCGCTTCTAGTGTCTCAACTTCTGTATCATC 879
QY 841 CTCTTGAAGCCCTCGATTAAGTTCCTGGAGAAGTGGTGCGCCAGATGCCCAATAACATTGAG 900
DB 880 CTCTTGAAGCCCTCGATTAAGTTCCTGGAGAAGTGGTGCGCCAGATGCCCAATAACATTGAG 939
QY 901 AAAAATTCAGCCGGGTGGCACTCTGGTGGTCTCTGAATTCAGTCAACCACTCTATGCT 960
DB 940 AAAAATTCAGCCGGGTGGCACTCTGGTGGTCTCTGAATTCAGTCAACCACTCTATGCT 999
QY 961 GGCATCAACTTCTCTTGTGCTGGTTCAGTTCAGTTGAGTTGGCAGACAGATCTCGTC 1020
DB 1000 GGCATCAACTTCTCTTGTGCTGGTTCAGTTCAGTTGAGTTGGCAGACAGATCTCGTC 1059
QY 1021 GACAAGGCGAGAACTGGGACATATGGCCCTGCACTAGTGTGAGGTTGGTAGAGAT 1080
DB 1060 GACAAGGCGAGAACTGGGACATATGGCCCTGCACTAGTGTGAGGTTGGTAGAGAT 1119
QY 1081 GTGATCATGCTCTTGGTCTTAAAGTTCCTTGGAGTGAAGTGTACTGAATTAAGTGTAT 1140
DB 1120 GTGATCATGCTCTTGGTCTTAAAGTTCCTTGGAGTGAAGTGTACTGAATTAAGTGTAT 1179
QY 1141 TCCTTGATTCGCTTCAGCTCAATTAATGCTTATCTGAATTCATTCAGTTCATGCTCCTT 1200
DB 1180 TCCTTGATTCGCTTCAGCTCAATTAATGCTTATCTGAATTCATTCAGTTCATGCTCCTT 1239
QY 1201 TTCTTCCAGTACTTGCATCCATTCGCTCACTCTTCACCCATAATGTAGTACACTACCTC 1260
DB 1240 TTCTTCCAGTACTTGCATCCATTCGCTCACTCTTCACCCATAATGTAGTACACTACCTC 1299
QY 1261 CATTTGTCTGCTGTCCAGCAGCACCCTCGGACAGGTTGAGAACTCAGAGCCACCTTTT 1320
DB 1300 CATTTGTCTGCTGTCCAGCAGCACCCTCGGACAGGTTGAGAACTCAGAGCCACCTTTT 1359
QY 1321 GAGACTGAAGCAAGCAAGTGTCTCTGA 1350
DB 1360 GAGACTGAAGCAAGCAAGTGTCTCTGA 1389

RESULT 3
US-09-768-781-5
; Sequence 5, Application US/09768781
; Patent No. US20020142376A1
; GENERAL INFORMATION:
; APPLICANT: MERKULOV, Gennady V. et al
; TITLE OF INVENTION: ISOLATED HUMAN TRANSPORTER PROTEINS,
; TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING HUMAN TRANSPORTER PROTEINS,
; TITLE OF INVENTION: AND USES THEREOF
; FILE REFERENCE: CL001057-CIP
; CURRENT APPLICATION NUMBER: US/09/768,781
; CURRENT FILING DATE: 2001-01-25
; NUMBER OF SEQ ID NOS: 7
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 5
; LENGTH: 17993
; TYPE: DNA
; ORGANISM: Human
; FEATURE:
; NAME/KEY: misc feature
; LOCATION: (1)-(17993)
; OTHER INFORMATION: n = A,T,C or G

US-09-768-781-5
Query Match 55.3%; Score 747.2; DB 10; Length 17993;
Best Local Similarity 97.1%; Pred. No. 2.1e-229;
Matches 761; Conservative 0; Mismatches 23; Indels 0; Gaps 0;
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QY 567 TGTGAGCCTGATCTCTGCAGAGGTTCCCTGGGTAGAGTTGTCTAATGGTATTTTCCCT 626
DB 15210 TGAATTCCTGTTTGTGTTTGTGTTTAAAGTTGCTAATGGTATTTTCCCT 15269
QY 627 GGTATCTGTACATATGGGGCCACCTTTGGCAATATGTTGGGTATCCAGATCAAGTACGA 686
DB 15270 GGTATCTGTACATATGGGGCCACCTTTGGCAATATGTTGGGTATCCAGATCAAGTACGA 15329
QY 687 TCAGTACAGATTCCTGCTTGGGCCACTAGAGTCTCTCATCACCATCTTGGCGGACATT 746
DB 15330 TCAGTACAGATTCCTGCTTGGGCCACTAGAGTCTCTCATCACCATCTTGGCGGACATT 15389
QY 747 GGAGATCACTTCCCGCTCCTGATTCCTGGTGTCTTCTCAGGCCACTTTGAAATTTGAAGGC 806
DB 15390 GGAGATCACTTCCCGCTCCTGATTCCTGGTGTCTTCTCAGGCCACTTTGAAATTTGAAGGC 15449
QY 807 TGTGCTTCTAGTGTCTCAACTTCTGTATCATCTCTTTGAGCCCTCGATTAAGTTCG 866
DB 15450 TGTGCTTCTAGTGTCTCAACTTCTGTATCATCTCTTTGAGCCCTCGATTAAGTTCG 15509
QY 867 GAGAGTGTGCGCCAGATGCCCAATAACATTGAGAAAAAATTCAGCCGGTGGCACTCT 926
DB 15510 GAGAGTGTGCGCCAGATGCCCAATAACATTGAGAAAAAATTCAGCCGGTGGCACTCT 15569
QY 927 GGTGTCTCTGATTTTCAGTCAACCATCTCTATGCTGCATCAACTTCTTCTGTGTGTCAGC 986
DB 15570 GGTGTCTCTGATTTTCAGTCAACCATCTCTATGCTGCATCAACTTCTTCTGTGTGTCAGC 15629
QY 987 TTTGAGTTGAGTTGGCAGACAGAGATCTCTCGACAAAGGGCAGAACTGGGGACATAT 1046
DB 15630 TTTGAGTTGAGTTGGCAGACAGAGATCTCTCGACAAAGGGCAGAACTGGGGACATAT 15689
QY 1047 GGGCTCTCACTATAGTGTGAGTTGGTAGAGATGTGATCATGGTCTTGGTTTAAAGTT 1106
DB 15690 GGGCTCTCACTATAGTGTGAGTTGGTAGAGATGTGATCATGGTCTTGGTTTAAAGTT 15749
QY 1107 CTTTGGAGTGAAGTGTGTAATTAATGCTCATTCCTTGAATTCCTTTCAGCTCAATTAT 1166
DB 15750 CTTTGGAGTGAAGTGTGTAATTAATGCTCATTCCTTGAATTCCTTTCAGCTCAATTAT 15809
QY 1167 TGTATCTGATTTCCATTCATGCTCTCTTTTCTTCCAGTACTTGCATCCATTGCG 1226
DB 15810 TGTATCTGATTTCCATTCATGCTCTCTTTTCTTCCAGTACTTGCATCCATTGCG 15869
QY 1227 CTCACCTTTCACCCATAATGTAGTACACTCCTCCATTGTGTCTGTCTGCTGTCACGACCC 1286
DB 15870 CTCACCTTTCACCCATAATGTAGTACACTCCTCCATTGTGTCTGTCTGCTGTCACGACCC 15929
QY 1287 TCGGACAGGTTGAGAACTCAGAGCCACCTTTTCAGACTGAAGCAAGGCAAGTGTGT 1346
DB 15930 TCGGACAGGTTGAGAACTCAGAGCCACCTTTTCAGACTGAAGCAAGGCAAGTGTGT 15989
QY 1347 CTGA 1350
DB 15990 CTGA 15993

RESULT 4
US-10-092-154-106
; Sequence 106, Application US/10092154
; Publication No. US20030054375A1
; GENERAL INFORMATION:
; APPLICANT: Rosen et al.
; TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
; FILE REFERENCE: PC009C1
; CURRENT APPLICATION NUMBER: US/10/092,154
; CURRENT FILING DATE: 2002-03-07
; NUMBER OF SEQ ID NOS: 2003
; Prior Application removed - See File Wrapper or Palm
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 106
; LENGTH: 531
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-092-154-106

Query Match 38.1%; Score 513.8; DB 9; Length 531;
Best Local Similarity 98.5%; Pred. No. 3.6e-155;
Matches 523; Conservative 5; Mismatches 2; Indels 1; Gaps 1;

Qy 747 GGAGATCACTTCCCGCCTCCTGATTTCTGGTCTCTTCAGCCACTTTGAAATTGAAGGC 806
Db 2 GGAGATCACTTCCCGCCTCCTGATTTCTGGTCTCTTCAGCCACTTTGAAATTGAAGGC 61
Qy 807 TGTGCCCTTCTAGTGTCAACTTCTGATCATCTCTTTGAGCCCTGATTAAGTTCTG 866
Db 62 TGTGCCCTTCTAGTGTCAACTTCTGATCATCTCTTTGAGCCCTGATTAAGTTCTG 121
Qy 867 GAGAAGTGGTGGCCAGATGCCCAATAAATTCAGCAAAATTTAGCCGGTGGGCACTCT 926
Db 122 GAGAAGTGGTGGCCAGATGCCCAATAAATTCAGCAAAATTTAGCCGGTGGGCACTCT 181
Qy 927 GGTGGTCTGATTTCAAGTCACTCTCTATGCTGGCATCAACTTCTCTGCTGTCAGC 986
Db 182 GGTGG-CTGATTTCAAGTCACTCTCTATGCTGGCATCAACTTCTCTGCTGTCAGC 240
Qy 987 TTTGCAGTTGAGTTGGCAGACAGAGATCTCGTCACAAAGGGCAGAACTGGGACATAT 1046
Db 241 TTTGCAGTTGAGTTGGCAGACAGAGATCTCGTYGACAAAGGGCAGAACTGGGACATAT 300
Qy 1047 GGGCCTGCACATATAGTGTGAGTTGGTAGAGAAATGTGATCATGTCTTGGTTTTAAGTT 1106
Db 301 GGGCCTGCACATATAGTGTGAGTTGGTAGAGAAATGTGATCATGTCTTGGTTTTAAGTT 360
Qy 1107 CTTTGGAGTGAAGTTTACTGAATTTACTGATCATGTCTTGGTTTTAAGTT 1166
Db 361 CTTTGGAGTGAAGTTTACTGAATTTACTGATCATGTCTTGGTTTTAAGTT 420
Qy 1167 TGTCTATCTGATTTCCATTTGCTTCTCTCTTCCAGTACTTGCATCCATTGG 1226
Db 421 TGTCTATCTGATTTCCATTTGCTTCTCTTCCAGTACTTGCATCCATTGG 480
Qy 1227 CTCACCTTCCACCCATATGATGATGACTACCTCCATTGTGTCTGTCGCA 1277
Db 481 CTCACCTTCCACCCATATGATGATGACTACCTCCATTGTGTCTGTCGCA 531

RESULT 5

US-09-764-847-106
; Sequence 106 Application US/09764847
; Patent No. US20020132767A1
; GENERAL INFORMATION:
; APPLICANT: Rosen et al.
; TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
; FILE REFERENCE: PC009
; CURRENT APPLICATION NUMBER: US/09/764,847
; CURRENT FILING DATE: 2001-01-17
; Prior application data removed - consult PALM or file wrapper
; NUMBER OF SEQ ID NOS: 2003
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 106
; LENGTH: 531
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-764-847-106

Query Match 38.1%; Score 513.8; DB 10; Length 531;
Best Local Similarity 98.5%; Pred. No. 3.6e-155;
Matches 523; Conservative 5; Mismatches 2; Indels 1; Gaps 1;

Qy 747 GGAGATCACTTCCCGCCTCCTGATTTCTGGTCTCTTCAGCCACTTTGAAATTGAAGGC 806
Db 2 GGAGATCACTTCCCGCCTCCTGATTTCTGGTCTCTTCAGCCACTTTGAAATTGAAGGC 61
Qy 807 TGTGCCCTTCTAGTGTCAACTTCTGATCATCTCTTTGAGCCCTGATTAAGTTCTG 866

Db 62 TGTGCCCTTCTAGTGTCAACTTCTGATCATCTCTTTGAGCCCTGATTAAGTTCTG 121
Qy 867 GAGAAGTGGTGGCCAGATGCCCAATAAATTCAGCAAAATTTAGCCGGTGGGCACTCT 926
Db 122 GAGAAGTGGTGGCCAGATGCCCAATAAATTCAGCAAAATTTAGCCGGTGGGCACTCT 181
Qy 927 GGTGGTCTGATTTCAAGTCACTCTCTATGCTGGCATCAACTTCTCTGCTGTCAGC 986
Db 182 GGTGG-CTGATTTCAAGTCACTCTCTATGCTGGCATCAACTTCTCTGCTGTCAGC 240
Qy 987 TTTGCAGTTGAGTTGGCAGACAGAGATCTCGTCACAAAGGGCAGAACTGGGACATAT 1046
Db 241 TTTGCAGTTGAGTTGGCAGACAGAGATCTCGTYGACAAAGGGCAGAACTGGGACATAT 300
Qy 1047 GGGCCTGCACATATAGTGTGAGTTGGTAGAGAAATGTGATCATGTCTTGGTTTTAAGTT 1106
Db 301 GGGCCTGCACATATAGTGTGAGTTGGTAGAGAAATGTGATCATGTCTTGGTTTTAAGTT 360
Qy 1107 CTTTGGAGTGAAGTTTACTGAATTTACTGATCATGTCTTGGTTTTAAGTT 1166
Db 361 CTTTGGAGTGAAGTTTACTGAATTTACTGATCATGTCTTGGTTTTAAGTT 420
Qy 1167 TGTCTATCTGATTTCCATTTGCTTCTCTTCCAGTACTTGCATCCATTGG 1226
Db 421 TGTCTATCTGATTTCCATTTGCTTCTCTTCCAGTACTTGCATCCATTGG 480
Qy 1227 CTCACCTTCCACCCATATGATGATGACTACCTCCATTGTGTCTGTCGCA 1277
Db 481 CTCACCTTCCACCCATATGATGATGACTACCTCCATTGTGTCTGTCGCA 531

RESULT 6

US-09-962-436-564
; Sequence 564 Application US/09962436
; Patent No. US20020081301A1
; GENERAL INFORMATION:
; APPLICANT: Sopdet, Daniel
; TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signal
; FILE REFERENCE: Sets
; CURRENT APPLICATION NUMBER: US/09/962,436
; CURRENT FILING DATE: 2001-09-25
; PRIOR APPLICATION NUMBER: US/60/235,082
; PRIOR FILING DATE: 2000-09-25
; PRIOR APPLICATION NUMBER: US/60/234,924
; PRIOR FILING DATE: 2000-09-25
; NUMBER OF SEQ ID NOS: 568
; SOFTWARE: PatentIn version 3.0
; SEQ ID NO 564
; LENGTH: 5096
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-962-436-564

Query Match 21.5%; Score 289.6; DB 10; Length 5096;
Best Local Similarity 54.6%; Pred. No. 3.8e-82;
Matches 627; Conservative 0; Mismatches 509; Indels 12; Gaps 2;

Qy 118 TCCACCTTTTGTACTGTGGGAGGCTGCATCTGCTTTGTACATGTTAGATCTATCGA 177
Db 110 TCCGTGTTCTGTTCTGTTGGCGGAGACAAACGGCGGCTCAGCCTGAGCAGCACCTACCGC 169
Qy 178 AAGAATAGTGAACCTTACCGGATGACATACACCTTTTCTTTTATGTTTTCATCCATT 237
Db 170 TCGGGGGGAGCCGATGTGCGAGCGCTGACGTGCTTTTCTCGCTACTGCTTGGCGG 229
Qy 238 ATGTTCCAGTTGACCTCAATTTTGTCCACAGAGATCTAGCCAAAGATAAACCCTATCA 297
Db 230 CTGTTGAGCTCAGCTTCTCTTGTACACCGGACCTCAGCGGACCGCCGCTCGTA 289
Qy 298 TTAATTTATGATCTAATCTCTTGGGACCTGTTATCAGATGTTTGGAGGCCATGATTAG 357

Db 2 ATTTCCCTGTTACAGTACTTATGGGCCATTTCGTCGAATATACCTGCGCATCCAGAT 61
 Qy 678 CAAGTAGATGACTACAGATTCCGCTTGGGCCACTAGAACTCTCTGTCATCACCATCTG 737
 Db 62 CAGCAATGATGATACCTACCAATTAAGCTACCCGCCGATAGAAATCTCTGTCGTGATGTG 121
 Qy 738 GCGGACATTTGAGATCACTTCCCGCTCTCTGATTCGTGCTCTCTCAGGCACCTTTGAA 797
 Db 122 CGGTTTTTGGAGGTTATCTCAGCTGAGTGAATCTGCGCATTTTTCATGATCTTGAA 181
 Qy 798 ATTGAAGGCTGTGCTTCTCTAGTCTCAACTTCCTGATCATCTCTTTTGAGCCCTGGAT 857
 Db 182 ACTGAAGAGCTACCCGCTTTTGTGTTAATCATATATTTTGTATCATTTTGTGGCACCCTGGCT 241
 Qy 858 TAAGTTCTGGAGAGTGGTGGCCAGATGCCCAATAACATTGAGAAAAATCTTCAGCCGGT 917
 Db 242 GGAGTTTTTGGAAAAAGTGGAGCTCATCTTCCTGGCAACAAAGAAAAATAATTCCAATATGGT 301
 Qy 918 CGGCACCTCTGGTGTCTGATTTTCAGTCAACCATCTCTATGCTGCGCATCAACTTCTCTTG 977
 Db 302 GGTACAGTACTGATGCTTTTCTTGATCACACTGCTATATGCTGCCATCAACTTCTCTG 361
 Qy 978 CTGCTCAGCTTTTGCAAGTTGAGTTGGCAGACAGAGATCTCGTCGACAAAGGCGAGAACTG 1037
 Db 362 CTGCTCAGCAGTGAACCTGTCAGTTGTGAGTGAACAAATAATTCAGGGGAGACAGAGGTG 421
 Qy 1038 GGGACATATGGCCCTGCACATATAGTGTGAGGTGGTAGAGATGTGATGCTGCTTGGT 1097
 Db 422 GGGCCATAGAATCTTACACTACAGCTTTTCAGTTTTTGTAGAAAAATGTGATAATGATTTGGT 481
 Qy 1098 TTTTAAGTCTTTTGGAGTGAAGTGTACTGAATTTACTGATCTCTGATTTGCTGCTTGA 1157
 Db 482 ATTTAGGTTCTTTGGAGGAAACTTTGCTGAAATTTGTGACTCATTAATTTGCCGTGCA 541
 Qy 1158 GCTCATTATTGCTTATCTGATTTCCATTGACTTTCATGCTCTCTTTTCTCCAGTACTGCA 1217
 Db 542 GTCATCATTAAGCTACCTATTTGGCCACTGGCTTTATGCTCTCTCTCTATCATGATTTGTA 601
 Qy 1218 TCATTGCGGTCA 1230
 Db 602 CCCATGGCAGTCA 614

RESULT 8

US-09-864-761-62
 ; Sequence 62, Application US/09864761
 ; Patent No. US20020048763A1
 ; GENERAL INFORMATION:
 ; APPLICANT: Penn, Sharon G.
 ; APPLICANT: Rank, David R.
 ; APPLICANT: Hanzel, David K.
 ; APPLICANT: Chen, Wensheng
 ; TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR
 ; TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
 ; FILE REFERENCE: Acomica-X-1
 ; CURRENT APPLICATION NUMBER: US/09/864, 761
 ; CURRENT FILING DATE: 2001-05-23
 ; PRIOR APPLICATION NUMBER: US 60/180,312
 ; PRIOR FILING DATE: 2000-02-04
 ; PRIOR APPLICATION NUMBER: US 60/207,456
 ; PRIOR FILING DATE: 2000-05-26
 ; PRIOR APPLICATION NUMBER: US 09/632,366
 ; PRIOR FILING DATE: 2000-08-03
 ; PRIOR APPLICATION NUMBER: GB 24263.6
 ; PRIOR FILING DATE: 2000-10-04
 ; PRIOR APPLICATION NUMBER: US 60/236,359
 ; PRIOR FILING DATE: 2000-09-27
 ; PRIOR APPLICATION NUMBER: PCT/US01/00666
 ; PRIOR FILING DATE: 2001-01-30
 ; PRIOR APPLICATION NUMBER: PCT/US01/00667
 ; PRIOR FILING DATE: 2001-01-30
 ; PRIOR APPLICATION NUMBER: PCT/US01/00664
 ; PRIOR FILING DATE: 2001-01-30

; PRIOR APPLICATION NUMBER: PCT/US01/00669
 ; PRIOR FILING DATE: 2001-01-30
 ; PRIOR APPLICATION NUMBER: PCT/US01/00665
 ; PRIOR FILING DATE: 2001-01-30
 ; PRIOR APPLICATION NUMBER: PCT/US01/00668
 ; PRIOR FILING DATE: 2001-01-30
 ; PRIOR APPLICATION NUMBER: PCT/US01/00663
 ; PRIOR FILING DATE: 2001-01-30
 ; PRIOR APPLICATION NUMBER: PCT/US01/00662
 ; PRIOR FILING DATE: 2001-01-30
 ; PRIOR APPLICATION NUMBER: PCT/US01/00661
 ; PRIOR FILING DATE: 2001-01-30
 ; PRIOR APPLICATION NUMBER: PCT/US01/00670
 ; PRIOR FILING DATE: 2001-01-30
 ; PRIOR APPLICATION NUMBER: US 60/234,687
 ; PRIOR FILING DATE: 2000-09-21
 ; PRIOR APPLICATION NUMBER: US 09/608,408
 ; PRIOR FILING DATE: 2000-06-30
 ; PRIOR APPLICATION NUMBER: US 09/774,203
 ; PRIOR FILING DATE: 2001-01-29
 ; NUMBER OF SEQ ID NOS: 49117
 ; SOFTWARE: Annonmax Sequence Listing Engine vers. 1.1
 ; SEQ ID NO 62
 ; LENGTH: 471
 ; TYPE: DNA
 ; ORGANISM: Homo sapiens
 ; FEATURE:
 ; OTHER INFORMATION: MAP TO AC005301.16
 ; OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 1.3
 ; OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 1.5
 ; OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 1.5
 ; OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 1.3
 ; OTHER INFORMATION: EXPRESSED IN HEL100, SIGNAL = 1.2
 ; OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 1.7
 ; OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 1.6
 ; OTHER INFORMATION: EXPRESSED IN BT474, SIGNAL = 1.1
 ; OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 1.3
 ; OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 1.8
 ; US-09-864-761-62

Query Match 13.1%; Score 176.8; DB 10; Length 471;
 Best Local Similarity 61.9%; Pred. No. 1.8e-46;
 Matches 280; Conservative 0; Mismatches 172; Indels 0; Gaps 0;

Qy 601 AGAGTTGTGCTAATGGTATTTTCCCTGCTATCTGTCACCTATGGGGCCACCCTTTGGCAAT 660
 Db 19 ATAGCATTGCTGATGACATTTTCCCTGTTATCAGTTACTTATGGGGCCATTGCGTSCAAT 78
 Qy 661 ATGTTGGCTATCCAGATCAAGTACGATGACTACAAAGATTTCGCTTGGGCCACTAGAATC 720
 Db 79 ATACTGGCCATCCAGATCAGCAATGATGATACCTACCAATAGCTACCCGCCGATGAATTC 138
 Qy 721 CTCGTCATCACCATCTGCGGCACATTTGGAGATCACTTCCCGCTCTCTGATTCGTGCTC 780
 Db 139 TTCTGTGCTGATGTGCGGTTTTTGGAGGTATCTCACTGTTAGTACTCTGCAATTT 198
 Qy 781 TTCAGCCACTTTGAAATTTGAAGCTGTGCGCTTCTCTAGTGTCTCAACTTCTGATCATC 840
 Db 199 TTCATTGTCATCTCTGAAACTGAAGAGCCTACCCGTTTTTGTAAATCATATATTTTGTATCA 258
 Qy 841 CTCCTTGAGCCCTGGATTAACTTCTGGAGAAAGTGTGCGCCAGATGCCCAATAACATTTAG 900
 Db 259 TGTGTGGCACCCTGGCTGGAGTTTTTGGAAAAAGTGAGGCTCATCTTCCTGGGCAACAAAGA 318
 Qy 901 AAAAATTCAGCCGGTGGCAGCTCTGCTGCTCTGATTTTCAGTCACCATCTCTCTATGCT 960
 Db 319 AATAATCCATATGTTGGGTACAGTACTGATGCTTTTCTTGATCACACTGCTATGCT 378
 Qy 961 GGCATCAACTTCTTCTGCTGCTGAGTTTGCAGTTTGGAGTTGGCAGACAGAGATCTCTGTC 1020
 Db 379 GCCATCAACTTCTCTCTGCTGCTGAGAGTGAAGTGAAGTGTGTCAGATGACAAAAATAATT 438
 Qy 1021 GACAAAGGGCAAACTGGGGACATATGGGCTT 1052


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; PRIOR APPLICATION NUMBER: PCT/US01/006668
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/006663
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/006662
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/006661
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/006670
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: US 60/234,687
; PRIOR FILING DATE: 2000-09-21
; PRIOR APPLICATION NUMBER: US 09/608,408
; PRIOR FILING DATE: 2000-06-30
; PRIOR APPLICATION NUMBER: US 09/774,203
; PRIOR FILING DATE: 2001-01-29
; NUMBER OF SEQ ID NOS: 49117
; SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
; SEQ ID NO 19197
; LENGTH: 294
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; OTHER INFORMATION: MAP TO AL121577.1
; OTHER INFORMATION: EXPRESSED IN BT474, SIGNAL = 1.2
; OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 2.3
; OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 1.6
; OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 1.8
; OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 2.4
; OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 1.5
; OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 2.2
; OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 2.2
; OTHER INFORMATION: EXPRESSED IN HEAL, SIGNAL = 2
; OTHER INFORMATION: EXPRESSED IN HBL100, SIGNAL = 3.3
; OTHER INFORMATION: NT HIT: g110835266, EVALUE 0.00e+00
; OTHER INFORMATION: EST HUMAN HIT: AI697050.1, EVALUE 0.00e+00
; OTHER INFORMATION: SWISSPROT HIT: P51811, EVALUE 5.00e-44
; OS-09-864-761-19197

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Query Match	5.4%	Score 73.2	DB 10	Length 294
Best Local Similarity	58.3%	Pred. No. 3.4e-13		
Matches 148	Conservative 0	Mismatches 103	Indels 3	Gaps
Qy	374	AGAAAGAGGAGCAGGAGGAGCCCTATGTCTCAGGCTCTACCCGAAAGA----	AGATGCTTAATAG	430
Db	28	AGTCAGGCAACATGAGAGCCTTATGTCTAGTATCACCAAGAGAGGCAATGCCAAAAA	87	
Qy	431	ATGGCGAGGAGGTGCTGTATAGAAATGGGAGGTGGGCCCACTCCATCCGACCCCTGGCTATGC	490	
Db	88	ATGGCTCTCTCAGAGGAGATTGAGAAGGAGGTGGGCCAGGAGCAAACTAATCACC	147	
Qy	491	ACGCAATGCTTACAAACATGTATGTCTACAGATCCAGCCTTCTCTGGGCTCAGTGC	550	
Db	148	ACCGATCAGCGTTTCAGCGGGGCTCGGTATCCAGCTTCTTTGGGCTCAGCCCCC	207	
Qy	551	TGACCTATCAGCTCTATGTGAGGCTGATCTCTGCAGAGGTTCCTCTGGGTAGAGTTG	610	
Db	208	TGACCTTACAGCTGTACATAAGTGTCTATGCAGCAGGACGTCACTGTTGGAAGAGTACGT	267	
Qy	611	TAATGGTATTTTCC	624	
Db	268	GTATTTTTTATTC	281	

RESULT 12

US-09-864-761-2467

; Sequence 2467, Application US/09864761

; Patent No. US20020048763A1

GENERAL INFORMATION:

; APPLICANT: Penn, Sharron G.

; APPLICANT: Rank, David R.

; APPLICANT: Hanzel, David K.

; APPLICANT: Chen, Wensheng

;	TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR
;	TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
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100	100

Query Match 4.8%; Score 65; DB 10; Length 477;
Best Local Similarity 62.2%;
Pred. No. 2.1e-10;
Matches 120; Conservative 0; Mismatches 70; Indels 3; Gaps 1;

[illegible]

Qy	491	ACCGCAATGCCTACAAACGTATGTCTACAGATCCAAAGCCTTCTCGGCTCAGTGC	550
Db	405	ACCGATCAGCGTTTCAACCGGGGTCGGTGATCCAGGCTTTCTTGGGCTCAGCCCCC	464
Qy	551	TGACCTATCAGCT	563
Db	465	TGACCTCTACAGCT	477

RESULT 13
US-09-864-761-10062/c
; Sequence 10062, Application US/09864761
; Patent No. US20020048763A1
; GENERAL INFORMATION:
; APPLICANT: Penn, Sharon G.
; APPLICANT: Rank, David R.
; APPLICANT: Hanzel, David K.
; APPLICANT: Chen, Wensheng
; TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR
; TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
; FILE REFERENCE: Aeomica-X-1
; CURRENT APPLICATION NUMBER: US/09/864,761
; CURRENT FILING DATE: 2001-05-23

Query Match	3.8%;	Score 51.6;	DB 10;	Length 486;
Best Local Similarity	56.5%;	Pred. No. 4.4e-06;		


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Matches 96; Conservative 0; Mismatches 74; Indels 0; Gaps 0;
QY 444 GCTGATGAATGGAGGTGGCCACTCATCGGACCCCTGGCTATGACCGAATGCCTA 503
Db 390 GATGCTGAAAGGAGATTGCAATTCCTAATCGGATAAATTTTCATGACGAGAAGGCTTT 331
QY 504 CAACGCTATGTCACAGATCCAAAGCTTCTCGGCTCAGTGGCCAGCTGACCTATCAGCT 563
Db 330 CAAGTACATGTCAGTGATTCAGGCTTTTCTCGGTTCTGTTCCACAATTAATTTTCAGAT 271
QY 564 CTATGTGAGCTGATCTCTGCAGAGTTTCCCTCGGTAGAGTTGTGCTAA 613
Db 270 GTATATCACTCTCACTATACGAGATGCGCTTTGAATAGAGTAAGTTGA 221

RESULT 14
US-09-764-877-853/c
; Sequence 853, Application US/09764877
; Patent No. US20020147140A1
; GENERAL INFORMATION:
; APPLICANT: Rosen et al.
; TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
; FILE REFERENCE: PC005
; CURRENT APPLICATION NUMBER: US/09/764,877
; CURRENT FILING DATE: 2001-01-17
; Prior application data removed - refer to PALM or file wrapper
; NUMBER OF SEQ ID NOS: 4031
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 853
; LENGTH: 832
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: SITE
; LOCATION: (824)
; OTHER INFORMATION: n equals a,t,g, or c
US-09-764-877-853

Query Match 3.2%; Score 43.4; DB 10; Length 832;
Best Local Similarity 49.8%; Pred. No. 0.0029;
Matches 110; Conservative 0; Mismatches 111; Indels 0; Gaps 0;
QY 1079 ATGTGATCATGTCTCTGTTTAAAGTTCTTTGGAGTAAAGTTACTGAATTAAGTCTG 1138
Db 500 AGGCTATCATTTCTAGGGAATTCGAGATGCTTGGAGTAAAGTGGAGAGCAGCTATG 441
QY 1139 ATTCCTTGATTCCTTGCAGCTCATTATTCTTATCTGATTTCCATTGACTTTCATGCTCC 1198
Db 440 TCACCTGTAATCAATCCCTTCTTCTCATTTGCTTAATCTTATCCAGATGTGATTCAACTCC 381
QY 1199 TTTTCTTCCAGTACTTGCATCCATTCGCTCACTTTCACCCATAATGTAGTAGACTACC 1258
Db 380 TTCACAGCTAACAAATTTTCCGAAGGTGTTCTCTCTGGGTGGATTTTCTGTTGGCCAGA 321
QY 1259 TCCATTGTCTGCTGTACACGACCCCTCGGACCAAGGTT 1299
Db 320 ATCCCTTGCTGTAGTCTTACAGTCTCCGAGGGATTCAGTT 280

RESULT 15
US-09-902-941-1883/c
; Sequence 1883, Application US/09902941
; Patent No. US20020172952A1
; GENERAL INFORMATION:
; APPLICANT: Henderson, Robert A.
; APPLICANT: Wang, Tonglong
; APPLICANT: Watanabe, Yoshihiro
; APPLICANT: Johnson, Jeffrey C.
; APPLICANT: Retter, Marc W.
; APPLICANT: Marnerakis, Margarita
; APPLICANT: Carter, Darrick
; APPLICANT: Fanger, Gary R.
; APPLICANT: Vedvick, Thomas S.
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; APPLICANT: Banqur, Chaitanya S.
; APPLICANT: McNabb, Andria
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
; TITLE OF INVENTION: AND DIAGNOSIS OF LUNG CANCER
; FILE REFERENCE: 210121.478C17
; CURRENT APPLICATION NUMBER: US/09/902,941
; CURRENT FILING DATE: 2001-07-10
; NUMBER OF SEQ ID NOS: 2002
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 1883
; LENGTH: 6799
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-902-941-1883

Query Match 2.9%; Score 38.8; DB 9; Length 6799;
Best Local Similarity 50.0%; Pred. No. 0.38; 97; Indels 0; Gaps 0;
Matches 97; Conservative 0; Mismatches 97; Indels 0; Gaps 0;
QY 342 GGAGGCCATGATTAAAGTACCTCACACTGTGGAAGAAAGAGGAGGAGGAGCCCTATGT 401
Db 859 GCGCATCTCTGGAGGAGAGCTGAAGCAGCTGGAGGAGAGTCCCGCAGAGAGGCGGA 800
QY 402 CAGCCTCACCCGAAAGAGATGCTAATAGATGGCGAGGAGTCTGATAGATGGGAGGT 461
Db 799 GCGTGTCAAGCTTGGAGCTGACGAGGTCAAGGAGAGCCTTGAAGAAAGCGCTGC 740
QY 462 GGGCAGCTCCATCCGACCCCTGGCTATGACCCGATGCTACACACGATGTCACAGAT 521
Db 739 GGGCGGAGTCAACCTTGGGCTGGCCATCGAGCCCAAGTCAGGACATCGAGTCCACAGT 680
QY 522 CCAAGCCTTCCTGG 535
Db 679 GTGTGTCCTTCTGG 666

Search completed: March 30, 2003, 06:55:09
Job time : 126.761 secs
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